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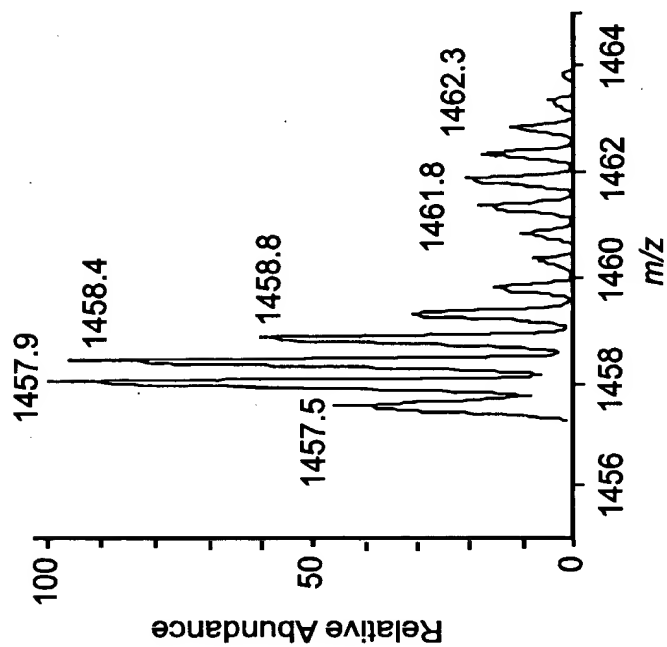


FIG. 1A

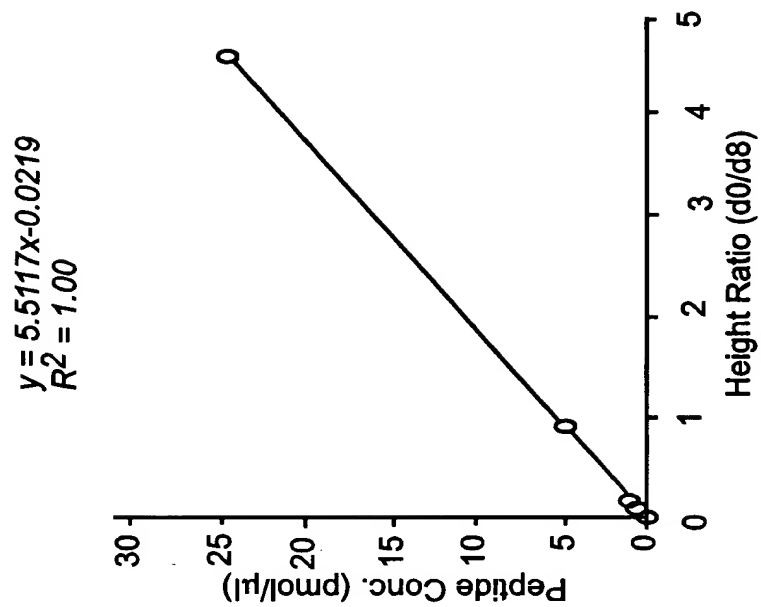


FIG. 1B

Mass spectrum of the peptide L-D-Q-W-L-C-E-K. The x-axis represents the mass-to-charge ratio (m/z) from 0 to 1400, and the y-axis represents Relative Abundance from 0 to 100. The base peak is at m/z 800, labeled $[M+2H]^{2+}$. The monoisotopic mass is 1522.7. Key fragments and transitions are labeled: K (approx. m/z 43), E (approx. m/z 283.2), C+biotin (approx. m/z 340), O (approx. m/z 357), W (approx. m/z 400), L (approx. m/z 573), Q (approx. m/z 1150), and D (approx. m/z 1237.7). A 'Biotin cleavage' label points to the transition between E and C+biotin.

4. 7. 20

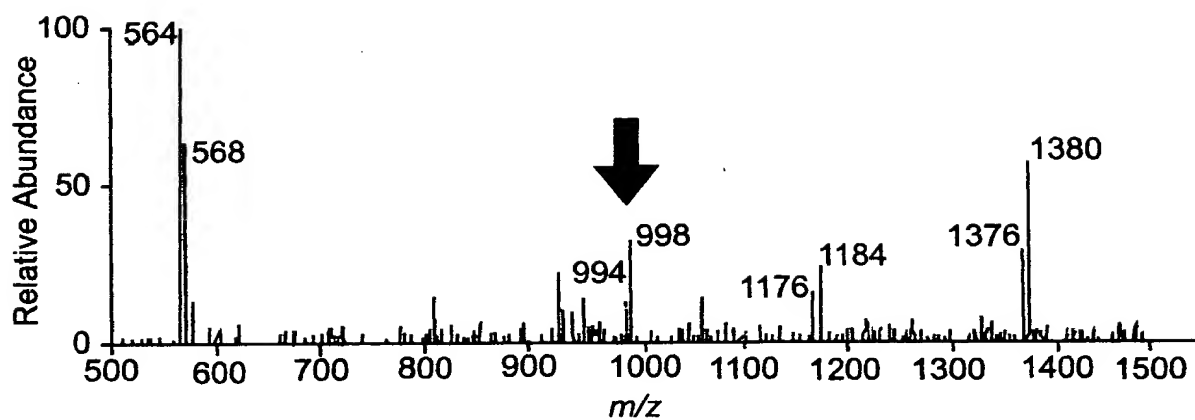


FIG. 3A

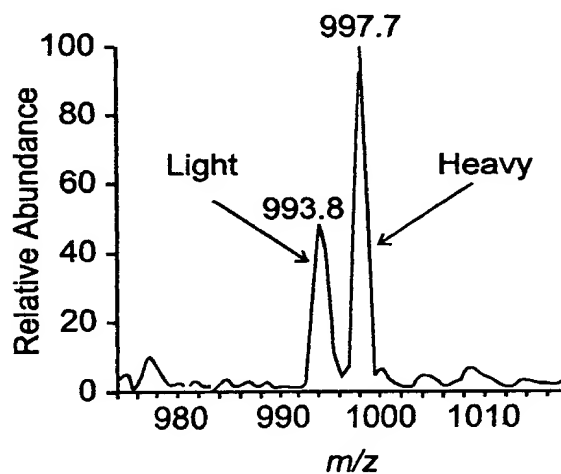


FIG. 3B

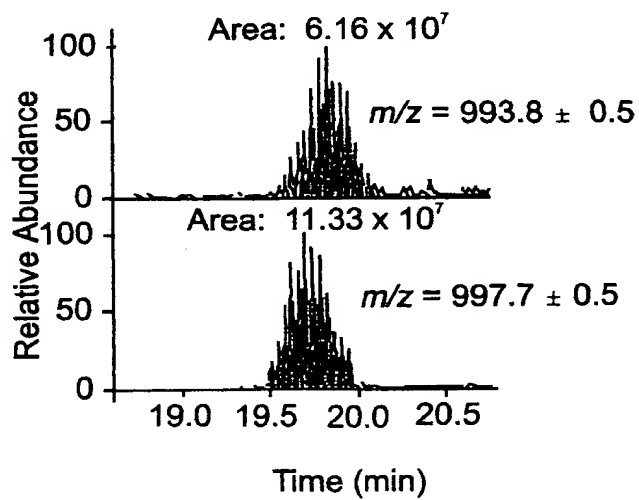


FIG. 3C

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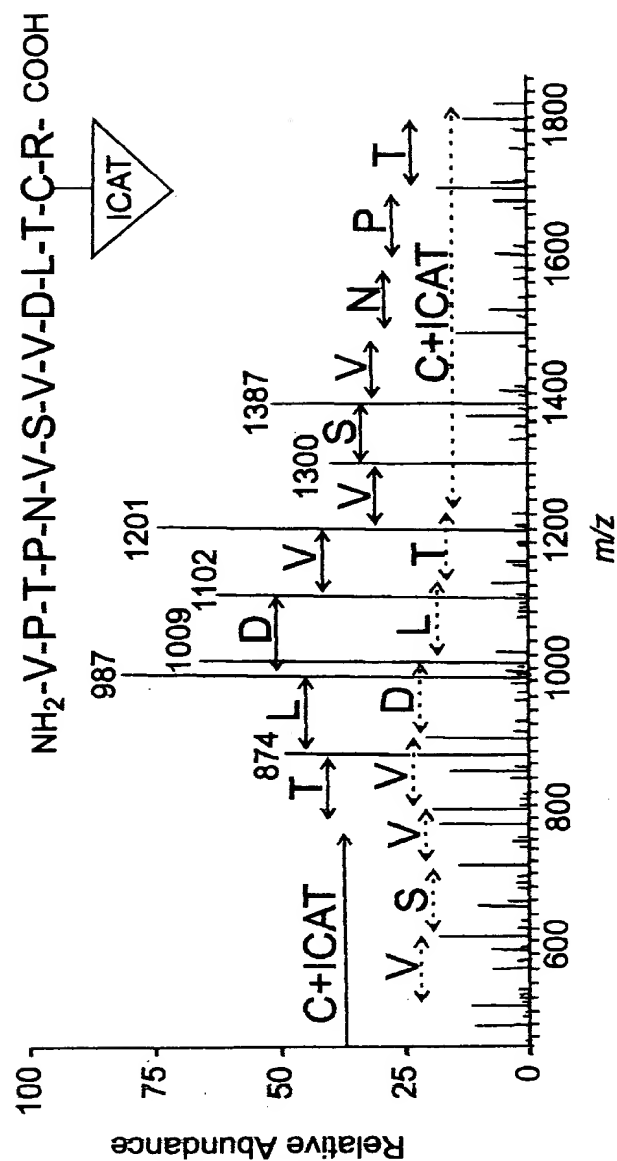


FIG. 4A

#	Rank/Sp	$(M+H)^+ \cdot 10^4$	Ions	Reference	Peptide
1.	1/ 1	1994.3 4.4675	17/26	G3P_RABIT	(R)VPTPNVSVVDLTC#R (SEQ ID NO:60)
2.	2/ 403	1995.1 2.7366	13/34	SLTRNGL	(E)LGKPPVLTANQVTIWEGLR (SEQ ID NO:61)
3.	3/ 3	1995.0 2.6591	16/36	FLP_LACCA	(N)LANPNVYTETLTAATVCTI (SEQ ID NO:62)
4.	4/ 209	1995.0 2.6335	14/36	A42912	(Y)LALLPSDAEGPHGQFVTDK (SEQ ID NO:63)
5.	5/ 381	1995.1 2.4634	13/38	H69373	(L)ALLVLVAPAMAAANGEDLRN (SEQ ID NO:64)

FIG. 4B

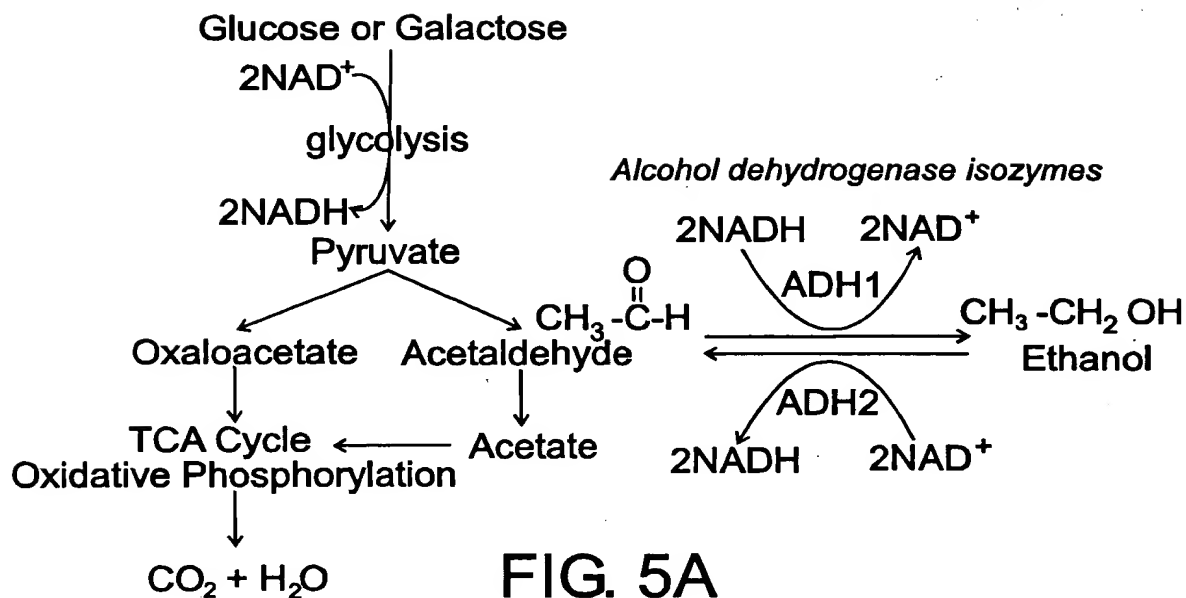
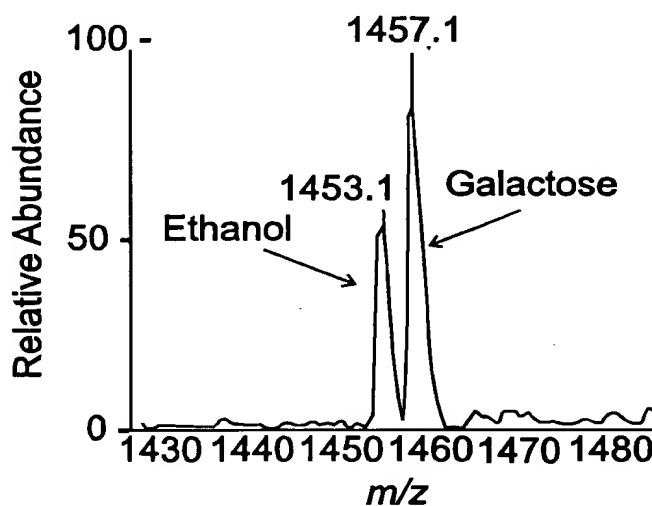


FIG. 5A

ADH1 : YSVC#HTDLHAWHGDWPLPVK

FIG. 5B



ADH2 : YSVC#HTDLHAWHGDWPLPIK

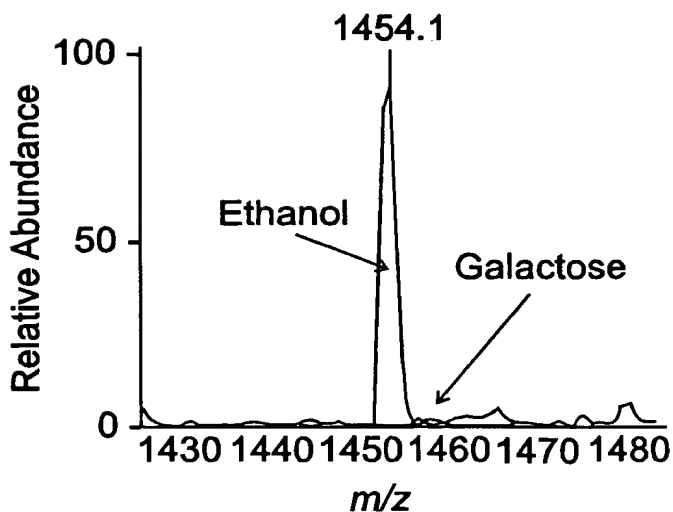


FIG. 5C

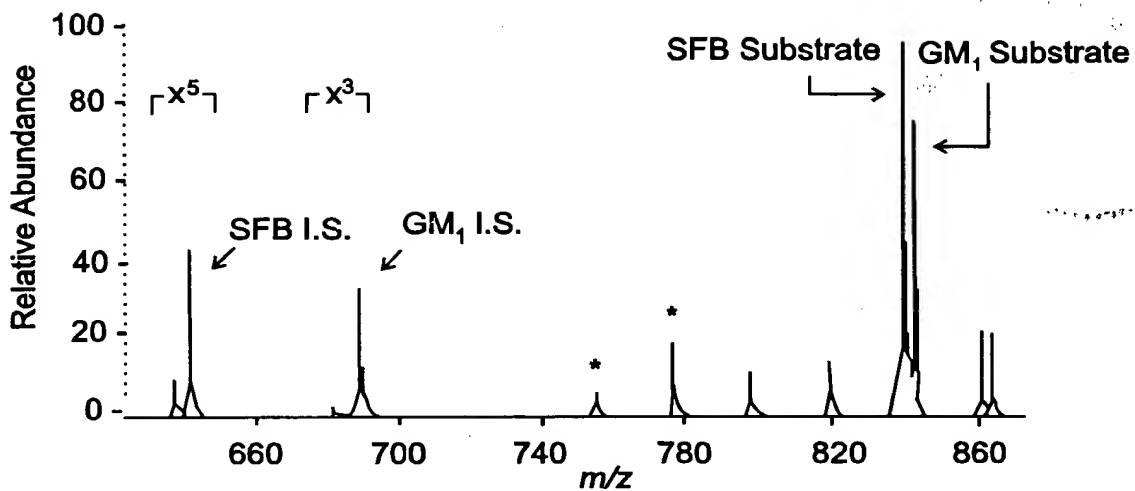


FIG. 6A

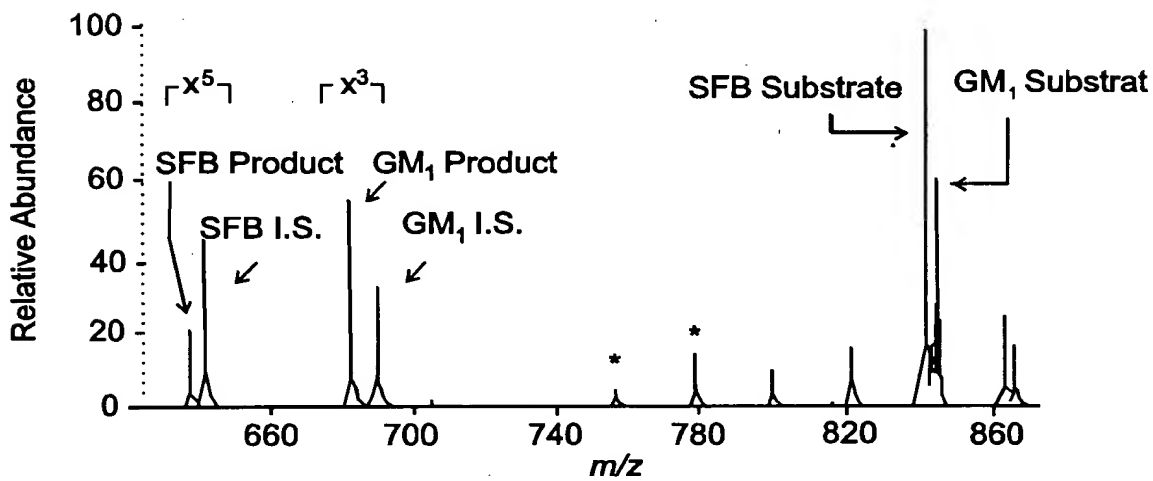


FIG. 6B

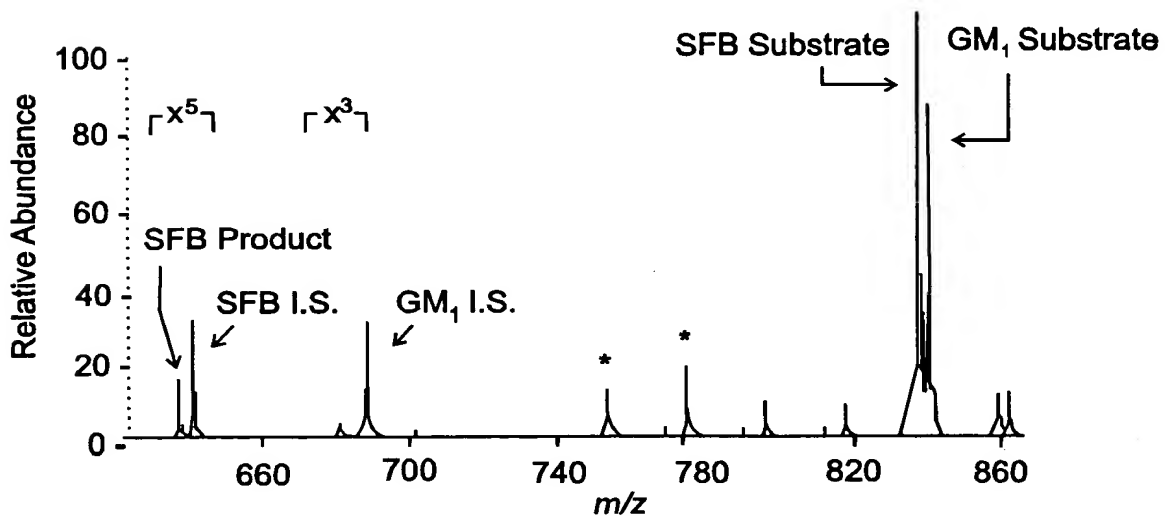


FIG. 6C

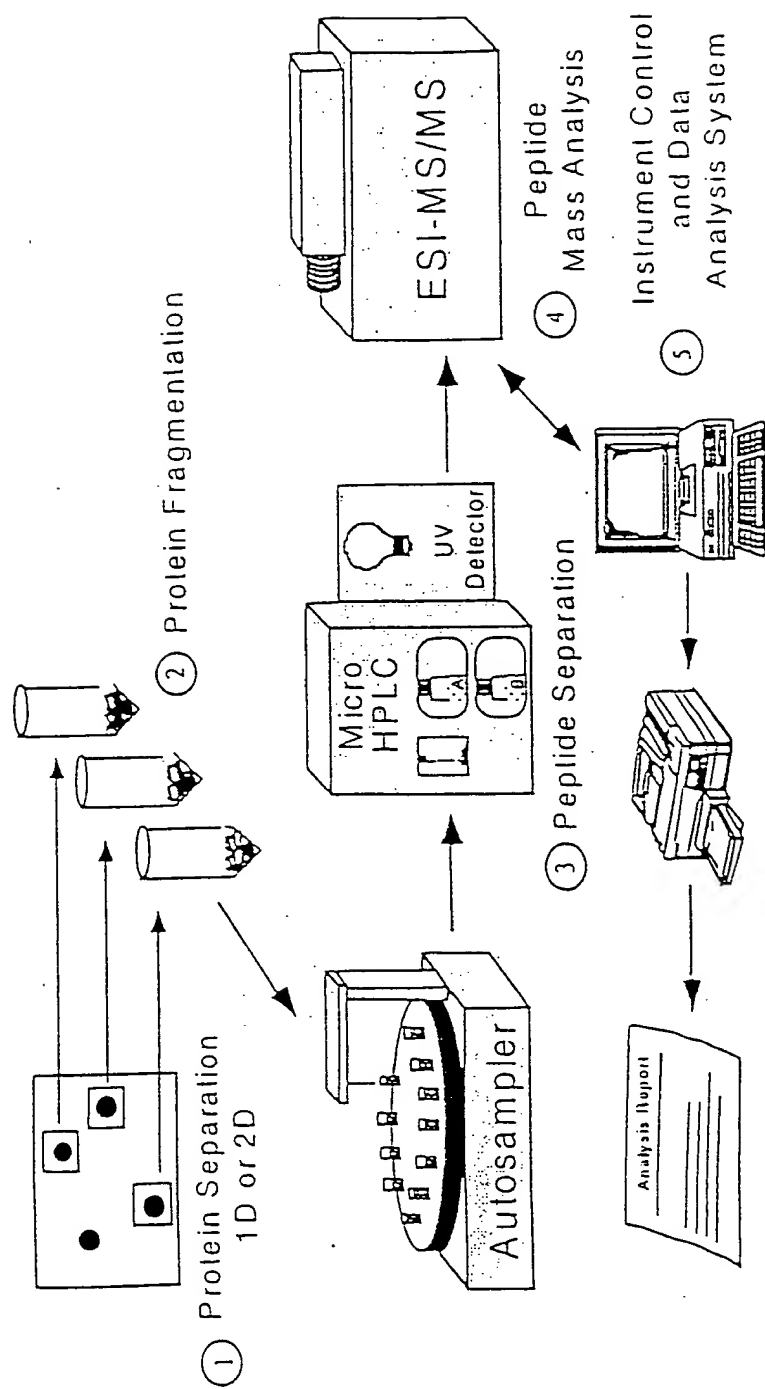


FIG. 7

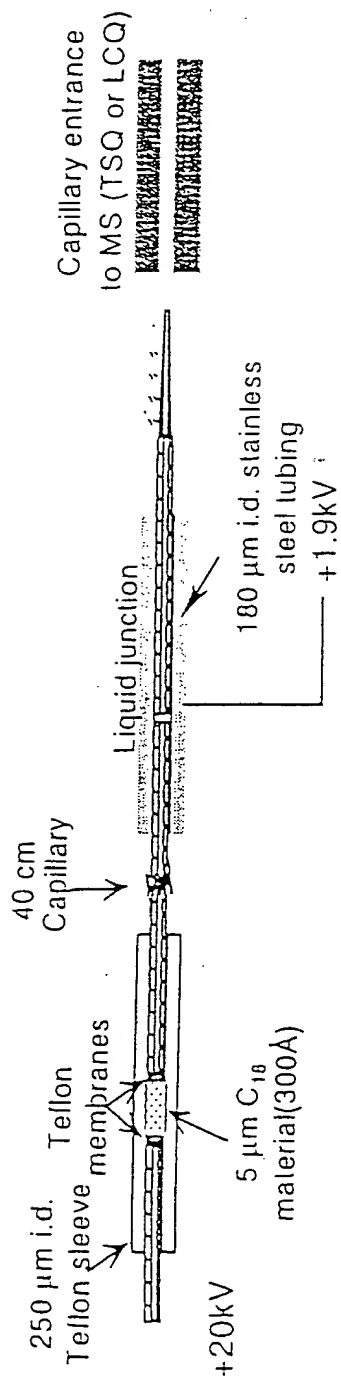


FIG. 8A

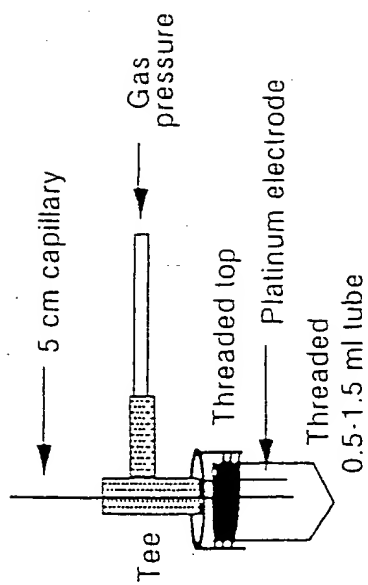


FIG. 8B